



# On the status of *Nomiapis valga* (Gerstäcker) and its distribution (Hymenoptera, Halictidae, Nomiinae), with an identification key for European *Nomiapis*

Thomas J. Wood<sup>1</sup>

1 Naturalis Biodiversity Center, Darwinweg 2, 2333 CR, Leiden, Netherlands

https://zoobank.org/6856018C-2E55-40EF-96BD-BA85362050F2

Corresponding author: Thomas J. Wood (thomas.wood@naturalis.nl)

Academic editor: Dominique Zimmermann ◆ Received 14 March 2025 ◆ Accepted 15 May 2025 ◆ Published 24 June 2025

## **Abstract**

Bees of the subfamily Nomiinae are renowned for their morphological diversity, particularly in the male sex, which often shows enormously modified hind legs and metasomal sterna. This has led to an array of generic and subgeneric classification systems, but nomiine bees also suffer from problems at the species level. This is well-illustrated in the genus *Nomiapis* Cockerell, 1919, which is particularly difficult to delineate in the female sex. One such case involves *Nomiapis valga* (Gerstäcker, 1872), described from southern Spain, which nominally has a fragmented distribution from Spain and southeastern Europe across to Central Asia, and for which type material is lost. New studies show that *N. valga* is actually entirely restricted to Spain, where it is strongly geographically separated from populations ranging from southeastern Europe to Central Asia; these latter populations are best referred to as *Nomiapis caucasica* (Friese, 1897) **sp. resurr.** Type material for *N. caucasica* is definitively located, and a neotype is designated for *N. valga* to fix the concept of this species. Additional support for the species status of *Nomiapis susannae* Arens, 2018, is presented, along with a modern identification key for European *Nomiapis* species. These results highlight the persistent problems with the alpha-level taxonomy of nomiine bees even within well-studied parts of their global range.

# Key Words

Cryptic species, lost type, museum collections, neotype, revisionary taxonomy

## Introduction

The subfamily Nomiinae contains around 620 species, with its center of diversity in the Afrotropics (Pauly 2009; Ascher and Pickering 2024; Bossert et al. 2025). Members of this subfamily show enormous variation in coloration (with the integument varying from dark to possessing iridescent metallic or enamel sections), pubescence, and morphology, with the male legs often grossly inflated and possessing pointed modifications and squamous hairs. This variation stands in strong contrast to the largest subfamily of Halictidae, Halictinae, most members of which have been described as "morphologically monotonous" (Michener 2007). Due to this morphological variation, traditional delineation of

nomiine genera is highly complex, with a lack of consensus in the literature (e.g., Warncke 1976; Pauly 1990, 2009, 2014; Pauly et al. 2001; Baker 2002; Michener 2007). Phylogenomic approaches are beginning to resolve these generic classification issues (Bossert et al. 2021; 2024; 2025), but many challenges remain.

In the West Palaearctic, nomine bees are represented by only a few lineages; in Europe, for example, only two genera (*Pseudapis* W.F. Kirby, 1900, and *Nomiapis* Cockerell, 1919) are present, with 11 species represented in total (Ghisbain et al. 2023). Although the overall diversity is low, recent studies have shown that West Palaearctic *Nomiapis* were more taxonomically complex than previously thought (Arens 2018; Wood and Le Divelec 2022), with the delineation of cryptic taxa

and description of new species. Due to the conspicuous variation present in male morphology (typically the legs and metasomal sterna) and the comparative lack of variation in females, identification of female specimens has caused numerous problems, leading to unrecognized diversity or incorrect geographic ranges based on female records only (Wood and Le Divelec 2022).

The two major modern revisions of the genus *Nomiapis* in its current form (e.g., Bossert et al. 2021) are Warncke (1976) and Baker (2002), after the more dated Friese (1897). Warncke treated *Nomiapis* as a subgenus within a broad genus, *Nomia* Latreille, 1804, and placed some taxa as subspecies of others, e.g., in the combination *Nomia femoralis* ssp. *valga* Gerstäcker, 1872. This revision was strongly criticized by Baker (2002) for its lack of precision, and while Baker's revision clarified the nomenclatural situation for several taxa, it did not fully deal with problems surrounding the *Nomiapis bispinosa* Brullé, 1833 species complex (Wood and Le Divelec 2022), and most pertinently for the present work, did not deal with *N. valga* in detail.

Nomiapis valga was described from southern Spain (Andalusia; Gerstäcker 1872), but the type material is lost (see results), and due to the scarcity of Spanish material in collections, the correctness of this locality has been questioned (Warncke 1976). Van der Zanden (1997) reported material from southern Spain, and Baker (2002) consequently thought said *locus typicus* to be reasonable, having never examined specimens himself. Warncke (1976) also considered the taxon N. caucasica (Friese 1897), which was described from the Caucasus, to be conspecific with N. valga (as a subspecies), although Baker (2002) treated N. caucasica as a synonym of N. equestris (Gerstäcker 1872). Impeding our understanding of the correctness of these two hypotheses is the fact that the type material of N. caucasica is currently considered lost (Warncke 1976; Baker 2002). A new study of museum collections, modern field collections, and rediscovery of type material in an unexpected location has allowed new light to be shed on these names and the overall distribution of the species N. valga.

# Methods

Some recently captured *Nomiapis* specimens were sent for genetic barcoding: a single midleg was removed from pinned specimens and sent to the Canadian Center for DNA Barcoding (CCDB) in Guelph, Canada, for DNA extraction and sequencing; specimens were sequenced following standardized high-throughput protocols (Ivanova et al. 2006). BeeCox1F1/BeeCox1R2 primers (Bleidorn and Henze 2021) were used to target the COI-5 region. All sequences are published on the Barcode of Life Database (BOLD) (https://portal.boldsystems.org/recordset/DS-NOMSYS).

Using the same primer, additional sequencing was conducted using the Oxford Nanopore GridION sequencer on two FLO-MIN114 flow cells with the SQK-NBD114.24 sequencing kit. The base calling was done with MinKNOW (v24.02.16), the run duration was set to 72 h, and super accuracy base calling was selected. The demultiplexing was performed with Guppy barcoder (v6.5.7). The consensus calling consisted of several steps combined together in a Snakemake (Mölder et al. 2021) pipeline: First, the reads (containing primers at both ends) were filtered by size (>=558, <=758) and quality ( $\geq 10$ ), and then reoriented with Cutadapt (v4.5, max error rate 20%, 80% coverage), which also removed flanking sequences. Then consensus sequences were generated using NGSpeciesID v0.3.0 (Sahlin et al. 2021) with Medaka polishing (v2.0.1). A final round of primer sequence trimming was performed with Cutadapt. Following this, multi-fasta files containing consensus sequences were written by using a custom script. Quality control and visualization of the processed FASTQ files were conducted using NanoPlot (De Coster et al. 2018) and MultiQC (Ewels et al. 2016).

Phylogenetic trees were supplemented with additional published sequences that were downloaded from Gen-Bank and the Barcode of Life Data System, as well as including sequences generated from historical specimens from the Naturalis Biodiversity Center collection as part of the Biodiversity Genomics Europe from specimens revised in the framework of this current work (https://portal.boldsystems.org/recordset/DS-BGEMS). Sequences were aligned using MAFFT (Katoh and Standley 2013). Aligned sequences were analyzed in Seaview (Gouy et al. 2010) using a maximum likelihood analysis, which was run with 1000 bootstraps. Intra- and interspecific distances were calculated using MEGA-X (Kumar et al. 2018). Genetic results are presented first before taxonomic changes are formalized.

Morphological terminology follows Michener (2007), with the exception of the marginal "zones" of the terga, which are referred to as marginal "areas." Unless otherwise stated, all specimens were examined, with determinations validated or corrected as necessary. An identification key is presented for males and females of *Nomiapis* species occurring in Europe to facilitate their identification. Photographs were taken using an Olympus E-M1 Mark II with a 60 mm macro lens. Additional close-ups were taken with the addition of a Mitutoyo M Plan Apo 10X infinity-corrected objective lens in combination with an Olympus M.Zuiko 2× teleconverter lens, a 10 mm Kenko DG extension tube, and a Meike MK-P-AF3B 10 mm extension tube. Photographs were stacked using Helicon Focus B (HeliconSoft, Ukraine), and plates were prepared in GNU Image Manipulation Program (GIMP) 2.10. Post-processing of some images was done in Photoshop Elements (Adobe Systems, USA) to improve lighting to highlight specific characters.

#### Abbreviations

**BBSL-USDA-ARS** Pollinating Insects Research Unit—

Dept. Biology, Utah State University, Logan, United States of America

NHMW Naturhistorisches Museum Wien,

Vienna, Austria

OÖLM Oberösterreichisches Landesmuse-

um, Linz, Austria

**OUMNH** Oxford University Museum of Natu-

ral History, Oxford, United Kingdom

RMNH Naturalis Biodiversity Center,

Leiden, the Netherlands

**TJWC** Personal collection of Thomas J.

Wood, Leiden, the Netherlands

**ZMHB** Museum für Naturkunde, Berlin,

Germany

## Results

### Genetics

Genetic sequences were available for nine European *Nomiapis* species, including the first for *N. caucasica*, *N. femoralis* (Pallas, 1773), *N. monstrosa* (Costa, 1861), and *N. susannae* Arens, 2018 (Fig. 1). The central result of Wood and Le Divelec (2022) was reproduced, with three well-supported species in the *bispinosa* group of species, which falls closest to *N. monstrosa*.

The sole sequence of N. valga presented by Wood and Le Divelec (2022) is now placed into context by sequences of N. femoralis (Czechia and Ukraine), N. caucasica (Greece), and additional sequences from southern Spain (Granada). Collectively, these sequences form a clade with bootstrap support of 95, which can be referred to as the *femoralis* group (*Nomia femoralis* sensu lato sensu Warncke). Individually, although only a small number of sequences were available, three distinct subclades were formed corresponding to N. femoralis (bootstrap support 82), N. valga (bootstrap support 86), and N. caucasica (bootstrap support 89). Nomiapis valga was separated from N. caucasica by an average genetic distance of 2.44% (range 2.43–2.49%) and from N. femoralis by an average genetic distance of 5.02% (range 4.54–5.47%). Nomiapis femoralis and N. caucasica were separated by an average genetic distance of 4.84% (range 4.55–5.13%). Consequently, three distinct clades were observed, and the genetic differences therefore support the morphological and biogeographical differences that are presented below.

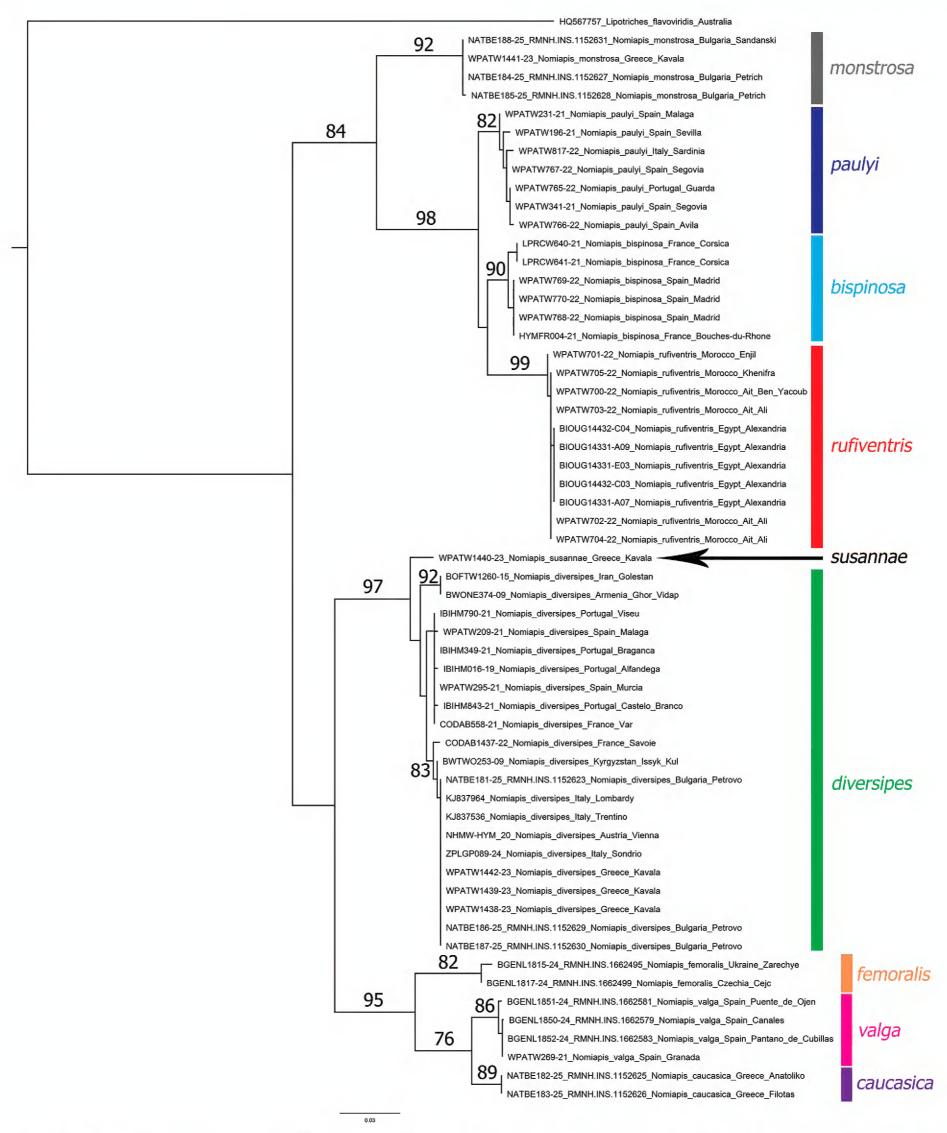
Finally, a large clade with bootstrap support of 97 was found containing *Nomiapis diversipes* (Latreille 1806) and *N. susannae*. *Nomiapis diversipes* sequences from Portugal in the west to Kyrgyzstan in the east formed a single monophyletic clade, with *N. susannae* placed as sister to this group. Within the broad *N. diversipes* 

clade, three subclades were found, namely 1) Armenia and northern Iran, 2) Portugal, Spain, and southern France, and 3) eastern France, Italy, Austria, Bulgaria, Greece, and Kyrgyzstan. *Nomiapis susannae* was separated from all N. diversipes sequences by an average genetic distance of 1.86% (range 1.75–2.14%). Within the broad N. diversipes clade, average intraspecific genetic distance was 0.68% (range 0.00–1.52%). Although the genetic distance between N. susannae and the broad N. diversipes clade is small in absolute terms, the minimum separation is greater than the maximum intraspecific distance inside N. diversipes, and the barcoded N. susannae specimen was captured in sympatry with N. diversipes specimens in Eastern Macedonia (around Kavala). The three *N. diversipes* from Kavala had identical sequences and were separated from N. susannae by 1.76%. In this context, combined with the observed morphological differences that matched the criteria of Arens (2018), it is appropriate to consider the two taxa distinct at the present time, though further study to clarify the overall range limits of N. susannae and identify the unknown female would be beneficial.

#### Nomiapis caucasica (Friese, 1897) sp. resurr.

*Nomia caucasica* Friese, 1897: 61, ♂ [Azerbaijan, NHMW, lectotype by present designation] (Fig. 2)

Material examined. Afghanistan • 1 ♂; Sarobi [Sarawbi]; 27 May 1951; Rolk leg.; OÖLM; Armenia • 1 ♂; pr. Eriwan [Yerevan], Parakar; 10 Jun. 1925; A. Schelkovnikow leg.; OÖLM; Azerbaijan • 3 &; Araxesthal [near Ordubad]; 1890; Reitter leg.; NHMW (lectotype by present designation); Greece • 1 ♀; Loutraki; 23 May 1964; M. Schwarz leg.; OÖLM • 1 ♀; Loutraki; 2 Jun. 1964; M. Schwarz leg.; OÖLM • 1 ♂; Loutraki, Golf von Korinth; 23–25 May 1962; M. Schwarz leg.; OOLM • 1 ♀; Rhodos, Archangelos; 2–14 Jun. 1996; M. Hradský leg.; OÖLM • 1  $\circlearrowleft$ ; Western Macedonia, Anatoliko, 6.3 km E, Church of the Holy Apostles; 1000 m a.s.l.; 15 Jun. 2024; T.J. Wood leg.; RMNH; RMNH.INS.1152625 • 1  $\varnothing$ ; Western Macedonia, Filotas, 3 km N of Antigonos; 600–700 m a.s.l.; 11 Jun. 2024; T.J. Wood leg.; RMNH; RMNH.INS.1152626; Iran • 1 ♂; E Iran, Mohammadabad; 1600 m a.s.l.; 3 May 1973; Exp. Nat. Mus. Praha; OÖLM • 1  $\delta$ ; Kuhanjan, SE Shiraz/Fars; 16 May 1978; K. Warncke leg.; OÖLM; ISRAEL • 1  $\circlearrowleft$ ; 2 km W Massada [Metsada]; 20 Apr. 1990; K. Warncke leg.; OÖLM • 4 ♂; Mizpe Ramon, 12 km W, Zisterne; 21 May 1991; K. Warncke leg.; OOLM • 4 3, 4 ♀; Mizpe Ramon, Borot Lotz; 950 m a.s.l.; 21 May 1991; K. Warncke leg.; OÖLM • 2 ♀; Wadi Abde; 13 May 1966; Bytinski-Salz leg.; OÖLM • 1 ♂; Wadi Ramon; 2 May 1966; Bytinskii-Salz leg.; OÖLM; Jordan • 1 ♀; west, Jordan Valley, S. Shuna; 25–26 Apr. 1996; Mi. Halada leg.; OÖLM; KAZAKHSTAN • 1 ♂, 1 ♀; centr., Lepsi, 6 km SE; 18 Jun. 1992; M. Halada leg.; OÖLM; TURKEY • 2 ♀;



**Figure 1.** Phylogenetic tree (maximum likelihood) of *Nomiapis* Cockerell, 1919 species based on the mitochondrial COI gene. *Lipotriches flavoviridis* (Cockerell, 1905) is used as an outgroup. Numbers adjacent to branches represent bootstrap support (values of < 75 are omitted).

Akseki, Antalya; 1300 m a.s.l.; 3 Aug. 1991; K. Warncke leg.; OÖLM • 1 ♂; Beysehir [Beyşehir]; 13–14 Jun. 1966; H.H.F. Hamann leg.; OÖLM • 1 ♂; Erzurum; 22 Aug. 1967; OÖLM • 1 ♀; Hakkari, Varegös/Mt. Sat; 1750 m a.s.l.; 6 Aug. 1986; K. Warncke leg.; OÖLM • 2 ♀; Malatya, 3 km NW of Darende, st. 126; 1400–1600 m a.s.l.; 31 Jul. 1983; H. v. Oorschot, H. v. d. Brink, H. Wiering leg.; RMNH;

RMNH.INS.1662589 • 1 \$\oplus\$; Malatya; 900 m a.s.l.; 27 Jun. 1984; K. Warncke leg.; OÖLM • 1 \$\oplus\$; Mut; 7 Jun. 1966; H.H.F. Hamann leg.; OÖLM • 1 \$\oplus\$; Mut; 26 May 1972; K. Warncke leg.; OÖLM • 1 \$\oplus\$; Nemrut Dağı, Karadut; 2 Jul. 1993; Mi. Halada leg.; OÖLM • 1 \$\oplus\$; Nevsehir [Nevşehir], road Göreme-Ürgüp, st. 132; 1000–1100 m a.s.l.; 8 Aug. 1983; H. v. Oorschot, H. v. d. Brink, H. Wiering leg.;



**Figure 2.** *Nomiapis caucasica* (Friese, 1897) lectotype male (NHMW). **A.** Label details; **B.** Habitus, profile view; **C.** Metasoma, dorsal view; **D.** Apical sterna, ventral view.

RMNH; RMNH.INS.1662591 • 1 ♀; Ulukışla; 19 Jun. 1973; K. Warncke leg.; OÖLM • 1 ♀; Yüksekova/Hakkari; 11 Aug. 1979; K. Warncke leg.; OÖLM; Turkmenistan • 1 ♂; Aschabad [Ashgabat]; Ahnger leg.; RMNH; RMNH. INS.1662593 • 2 ♂; Aschabat, 40 km W Firyuza; 6 Jun. 1993; M. Halada leg.; OÖLM • 1 ♀; Кара-Кала, Туркм. [Kara-Kala, Turkmenistan]; 11 Jun. 1953; Крыжановский [Kryzhanovsky] leg.; OÖLM; Uzbekistan • 1 ♂; Samarkand - Aman Kutan; 1 Jun. 1919; RMNH; RMNH. INS.1662592; West Bank • 1 ♂; Jericho; 3 Apr. 1909; F. Morice leg.; OUMNH • 5 ♂; Jericho, Wadi Qilt; 21 Apr. 1990; K. Warncke leg.; OÖLM • 1 ♂; Shelomzziyon war memorial; 12 Apr. 2000; S.P.M. Roberts leg.; TJWC.

**Notes.** The location of the type material of *N. caucasica* has been a mystery until now. Friese (1897: 62) wrote, "Im Museum Wien befinden sich 4 ♂ aus dem Kaukasus-Gebeit (Arexesthal 1890)." For inexplicable reasons, Warncke (1976: 113) wrote, "*N. caucasica* FRIESE, 1922, ♂, Kaukasus, Typus. Mus. Berlin, ist *N. femoralis* ssp. *valga!* In seiner Beschreibung kennzeichnete FRIESE dagegen *N. equestris!* Beschreibung und Typus (sowie Paratypen) stimmen nicht überein! Syn. n.!". The "type" specimen in the ZMHB collection is actually a specimen from Sarachs in Turkmenistan, which was identified by Friese (R. Le Divelec, *pers. comm.*), but is clearly of no

type value as it does not match the collecting information in the original publication.

That the type of *N. caucasica* clearly should not be in the ZMHB was noted by Baker, who correctly stated that it should be found in the NHMW based on the original publication and who then listed *N. caucasica* as a synonym of *N. equestris*. This synonymy is based on the fact that Friese's description of *N. caucasica* is partially copied from Gerstäcker's description of *N. equestris*, but Baker did not inspect the type material of *N. caucasica* either, and so this synonymy is speculative.

Searches in the NHMW collection in 2024 were not able to retrieve any specimens of N. caucasica (D. Zimmermann and R. Le Divelec,  $pers.\ comm$ .), but a loan form dated 2 April 1965 was uncovered, this detailing a loan of 38 nomiine specimens to M.A. Diniz at the University of Coimbra (Portugal), including "Nomia caucasica 3 1 Ex." Searches at the University of Coimbra museum collection as well as communication with Diniz were unable to locate these specimens, including N. caucasica (H. Gaspar,  $pers.\ comm$ .). This explains why one of the N. caucasica specimens was missing from the NHMW collection, but what of the other three?

During searches of the Warncke collection (OÖLM), three specimens were found that had been determined by

Warncke as "Nomia femoralis ssp. valga." These had the collecting information "Araxesthal, Reitter 1890" and were labeled by Friese as "Nomia caucasica." Warncke had even added "Para-type" labels, raising the point that he was clearly aware of their type status (Fig. 2A). These three specimens are indisputably the remaining syntypes of *N. caucasica* that had been taken from NHMW by Warncke at an unknown date due to their collecting labels and consistency with Friese's original description. One of them is designated as the lectotype, the type series now having been returned to the NHMW. It is unclear if Warncke took these from the NHMW collection before or after his 1976 publication.

Now that the type material has been definitively located, its identity can be established and the hypotheses of Warncke (conspecific with *valga*) and Baker (conspecific with *equestris*) can be assessed. The specimen is clearly not conspecific with *N. equestris*, as the apical margin of S4 has a deep semi-circular excavation, and the apical margin of S5 is weakly emarginate with two pincer-like, slightly bent digitiform projections (Fig. 2D; the narrowly separated, raised, convex, tongue-like processes of Baker 2002: 57). Moreover, the apical process of the hind tibia is flattened, more-or-less rectangular, and with the apical margin truncate (Fig. 2B). As *N. equestris* males have the hind margin of S4 narrowly incised, S5 with a large circular excavation, and the apical process of the hind tibia narrow and sharply pointed, Baker's hypothesis can be rejected.

Separation from *N. valga* is more challenging. As the lateral margins flanking the emargination of S4 are produced into low-lying posteriorly projecting ridges, it can be separated from *N. femoralis*, which has the lateral margins produced elevated projections with triangular apexes, Warncke's combination of *N. femoralis* ssp. *valga* being rejected by most subsequent workers (Ebmer 1988; Baker 2002; Astafurova and Pesenko 2006). Consultation of a series of specimens from Spain (*N. valga*) and eastern specimens (*N. caucasica*) has allowed discriminative characters to be identified in the male sex:

- 1. Density of punctures on the disc of T2
- 2. Punctation of the marginal areas of T2–3
- 3. Tergal sculpture, including sculpture of marginal areas

The lectotype of *N. caucasica* displays 1) relatively sparse punctures on the disc of T2, with two shining areas mediolaterally that clearly exceed the diameter of a puncture in size (Fig. 2C); 2) the marginal areas of T2–3 have a very narrow row of punctures at their base, these punctures covering <½ of the length of the marginal area (Fig. 2C); and 3) the terga have relatively weak microsculpture sculpture and are broadly shining, particularly the marginal areas (Fig. 2C).

In contrast, specimens from Spain display 1) relatively dense punctures on the disc of T2, without impunctate shining areas mediolaterally that clearly exceed the diameter of a puncture in size, at most with slight hints of such an area (Fig. 3C); 2) the marginal areas of T2–3 have 3–4

rows of punctures at their base, these occupying the basal  $\frac{1}{3}-\frac{1}{5}$  of the marginal areas (Fig. 3C); and 3) the terga have slightly stronger microsculpture and are comparatively more dull, particularly on the marginal areas, which are never polished and shining (Fig. 3C). Finally, the lectotype of N. caucasica has the scutellum laterally rounded, whereas specimens of N. valga from Spain have the scutellum laterally mucronate, with posteriorly projecting teeth. Baker (2002: 57, couplet 26) considered males of N. valga to have the scutellum laterally without posteriorly projecting spines, but he never examined specimens from Spain, which show the spines consistently based on the limited number of specimens examined to date. Specimens from the east are variable, sometimes showing spines and sometimes not; it therefore does not seem to be a stable character in eastern populations, even though it may be stable (always present) in N. valga. Overall, these characters taken in combination allow the consistent separation of male N. caucasica and N. valga regardless of geographic origin, and N. caucasica sp. resurr. is returned to species status.

As the type material of *N. valga* is lost, with Friese, Warncke, and Baker unable to locate it in the ZMHB, along with modern searchers (R. Le Divelec, pers. comm.), and the clear taxonomic complexity in separating N. valga and N. caucasica, it is desirable to designate a neotype for N. valga. In line with the conditions of article 75.3 (ICZN 1999), this neotype (Fig. 3) is needed to decisively settle the identity of N. valga, preserving nomenclatural stability for Iberian populations. The selected specimen is from the province of Granada, which falls within Andalusia, the stated terra typica of Gerstäcker. The selected specimen is male, matching the sex in which the species was originally described. This neotype selection allows the name N. valga to continue to be applied to Spanish populations. The neotype is deposited in the RMNH collection (unique reference number: RMNH.INS.1662584).

Concerning the distribution of *N. caucasica*, due to historical problems with the species concepts, it is necessary to clarify some points here. Astafurova and Pesenko (2006) list N. caucasica as a synonym of N. valga following Warncke and give a broad distribution from Spain to Pakistan. This also includes Czechia; this record is outside of the range of N. valga sensu Warncke and derives from Van der Zanden (1997). Van der Zanden (1997: 756) reported two males of "Pseudapis valga" from Czechia (Čejč), one male from Spain (Granada), one male from Turkmenistan (Ashgabat), and one male from Uzbekistan (Samarkand; reported as collected in 1959 but in reality collected in 1919 based on the label). All specimens are now part of the RMNH collection and were examined; the two males from Czechia are N. femoralis, the male from Spain is N. valga, and the two males from Turkmenistan and Uzbekistan are N. caucasica (full specimen details are found in their respective species entries). This inconsistent identification is unfortunately characteristic of the later work of Van der Zanden (see Baker 2002; Wood and Le Divelec 2022), but the re-examination of material allows for these literature references to be clarified.



**Figure 3.** *Nomiapis valga* (Gerstäcker, 1872) neotype male (RMNH). **A.** Label details; **B.** Habitus, profile view; **C.** Metasoma, dorsal view; **D.** Hind leg and metasoma, profile view.

Finally, Warncke (1980: 370) reported *N. femoralis* ssp. *valga* from Iran and Afghanistan; specimens from these two records were examined in the OÖLM collection, with them corresponding to *N. caucasica*. Overall, *N. caucasica* can be seen as the eastern counterpart of *N. valga*, which is here restricted in distribution to just Spain (Fig. 4).

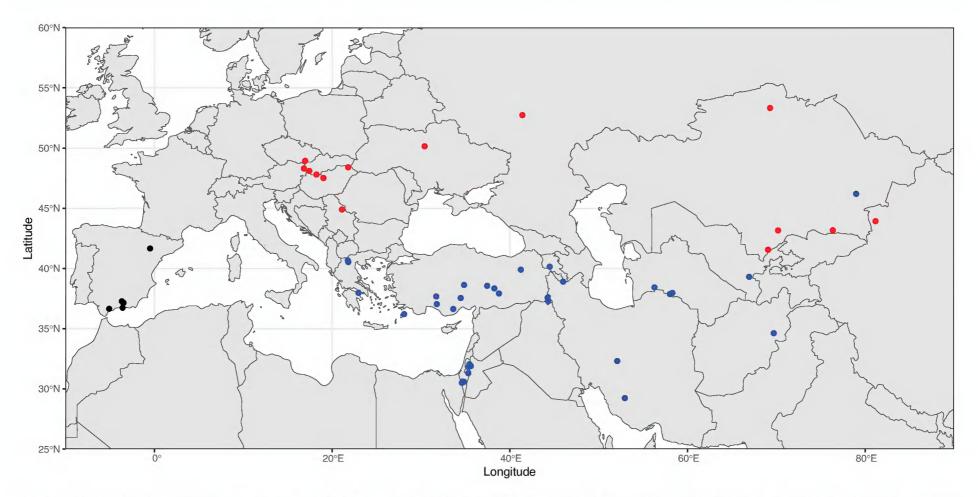
**Distribution.** North Macedonia, Greece (including Samos, Lesbos, and Rhodos), Turkey, Cyprus, Israel and the West Bank, Jordan, Armenia, Azerbaijan, Iran, Kazakhstan, Turkmenistan, Uzbekistan, Tajikistan, Afghanistan, and Pakistan (Warncke 1976; 1980 *partim*, as *Nomia femoralis* ssp. *valga*; Baker 2002 *partim*, as *Nomiapis valga*; Astafurova and Pesenko 2006 *partim*, as *N. valga*; Ebmer 2014 *partim*, as *Pseudapis valga*; Pauly 2015 *partim*, as *N. valga*) (Fig. 4).

#### Nomiapis femoralis (Pallas, 1773)

Apis femoralis Pallas, 1773: 731, ♂ [Kazakhstan, ZMHB, not examined]

Material examined. Austria • 2 ♂; N.O. Marchfeld, Oberweiden (Sandberge); 7 Jul. 1954; F. Koller leg.; A.W. Ebmer det. 1974; OÖLM • 3 ♂; N.O. Marchfeld, Oberweiden (Sandberge); 21–22 Jul. 1954; F. Koller

leg.; A.W. Ebmer det. 1974; OÖLM • 1 ♂, 1 ♀; Ober Weiden, A. i. [Austria Inferior]; [undated]; Mader leg.; M. Schwarz det. 1986; OÖLM • 1 ♂; Oberweiden; 27 Jun. 1931; J. Kloiber leg.; A.W. Ebmer det. 1974; OÖLM • 1 ♀; Oberweiden; 28 Aug. 1949; M. Kocourek leg.; M. Kocourek det.; OÖLM; China • 1 ♂, 1 ♀; Boro Horo mts. [Xinjiang], Jinig [Yining/Gulja], Ining-H-Sein; 27 Jul. 1991; M. Snížek leg.; A. Pauly det. 2008; OÖLM; Czechia • 1 ♀; Čejč - Mor. [Moravia]; 16 Jul. 1939; O. Sustera leg.; RMNH; RMNH.INS.1662494 • 1 ♂; Čejč - Mor. [Moravia]; 27 Jul. 1939; O. Sustera leg.; RMNH; RMNH.INS.1662500 • 1 ♂; Čejč - Mor. [Moravia]; 27 Jul. 1939; V. Zavadil leg.; K. Warncke det.; OÖLM • 1 &; Čejč - Mor. [Moravia]; 1 Aug. 1939; O. Sustera leg.; RMNH; RMNH.INS.1662499 • 1 3, 1 ♀; Čejč - Mor. [Moravia]; 27 Jul. 1941; V. Zavadil leg.; K. Warncke det.; OÖLM • 1 ♀; Čejč - Mor. [Moravia]; 1–31 Aug. 1958; M. Kocourek leg.; K. Warncke det.; OÖLM; Hungary • 1 ♀; Budapest; 19 Jun. 1927; E. Stöckhert leg.; K. Warncke det.; OÖLM • 1 ♀; Budapest; [undated]; RMNH; RMNH.INS.1662493; KAZAKHstan • 1 3; Alma Ata, 30 km W, Aksaj; 16 Jul. 1981; M. Kocourek leg.; M. Kocourek det.; OÖLM • 1 ♀; Darbaza, 40 km N Tachkent; 30 May 1994; Ma. Halada leg.; A. Pauly det.; OÖLM • 1 ♀; Džambul [=Taraz] env,



**Figure 4.** Distribution map based on specimens examined during this study for *Nomiapis valga* (Gerstäcker, 1872) (black circles), *Nomiapis femoralis* (Pallas, 1773) (red circles), and *Nomiapis caucasica* (Friese, 1897) (blue circles). Distributional data presented here are not comprehensive, and additional references in the main text should also be consulted in the context of the revisions to *Nomiapis* species concepts presented here.

Kara Tau, průsmyk [pass] Ujuk; 1000 m a.s.l.; 3 Jun. 1980; Z. Pádr leg.; K. Warncke det.; OÖLM • 1  $\stackrel{\wedge}{\circ}$ , 1 ♀; Боровск [Borovsk= Borovoye= Burabay], лес тех Кокчет [Kokchetav] р. Акмол [Akmol]; 2–5 Jul. 1932; В. Попов [V. Popov] leg.; K. Warncke det.; OÖLM; Russia • 1  $\circlearrowleft$ ; Tambow [Tambov]; Ф. Моравица [F. Morawitz] leg.; K. Warncke det.; OÖLM; SERBIA • 1 ♂; Deliblatska Peščara; 2 Aug. 1958; RMNH; RMNH. INS.1662498 • 1 ♂; Deliblatska Pescara [Deliblatska Peščara], Banat - Yoego Slavië; 12 Jul. 1950; Adamovic leg.; RMNH; RMNH.INS.1662497; SLOVAKIA • 1 ♂; Čenkov [Čenkovce]; 1-31 Jul. 1964; M. Kocourek leg.; M. Kocourek det.; OÖLM • 1 ♀; Chotin; 22 Jul. 1962; Z. Pádr leg.; OÖLM • 1 ♂; Chotin; 1–31 Aug. 1962; M. Kocourek leg.; K. Warncke det.; OÖLM • 3 ♂; Somotor; 1-31 Jul. 1952; M. Kocourek leg.; M. Kocourek det.; OÖLM; Ukraine • 1 ♂; Киевская Обл [Kiev Oblast], Каневский р-н, о-в Заречье [Zarechye/Zarichchya]; 20 Jun. 1950; A. Осычнюк [A. Osytshnjuk] leg.; RMNH; RMNH.INS.1662495 • 1 ♀; Киевская Обл [Kiev Oblast], Каневский р-н, о-в Заречье [Zarechye/Zarichchya]; 23 Jun. 1950; А. Осычнюк [A. Osytshnjuk] leg.; RMNH; RMNH.INS.1662496.

**Notes.** The *locus typicus* is "*in deserto ad Iaïkum*," which Astafurova and Pesenko (2006: 78) give as "in a desert on the bank of the Ural River, western Kazakhstan." Revision of material supports the distribution of this species given by Pauly (2015) with the exception of a record from Rhodes; this specimen was determined as "*valga/femoralis* female" by Pauly in 2008 (direct information from the determination label) but seems to have been accidentally encoded and displayed as *N. femoralis*; it is actually referable to *N. caucasica* (see material examined above). *Nomiapis femoralis* is clearly a species of Eurasian steppe

and does not descend into the southern Balkans and Turkey, where *N. caucasica* can be found (Fig. 4).

The record from Spain (Barcelona) given by Warncke remains a mystery, as no Spanish specimens in the OÖLM collection could be found. Wood et al. (2020) were able to inspect via photograph (Fig. 5) one of the specimens housed in the BBSL-USDA-ARS collection that was determined to be *Nomiapis monstrosa* but which cannot belong to this species, as the male lacks a digitiform process at the base of the hind tibia. The specimens are of the valga/caucasica type due to the sternal morphology (i.e., with curved processes found medially on the 4<sup>th</sup> sternum), and given the geographic location in Spain, it is considered much more likely that they represent N. valga rather than N. caucasica, whose closest populations would be in the southern Balkans, some 1,500 km distant. The position is therefore taken that Warncke's (1976) record from Barcelona represents N. valga, and N. femoralis is excluded from the Spanish fauna.

Finally, based on the limited material examined here, it is important to highlight the clear decline of *N. femoralis* in Europe. Across multiple Central European countries, *N. femoralis* was last recorded in Austria in 1954 (see also Kratschmer et al. 2021), Czechia in 1958 (regionally extinct, Hejda et al. 2017), Serbia in 1958, and Slovenia in 1964. Although the species persists in Hungary, with specimens collected in 2024 (D. Zimmermann, *pers. comm.*), this fits the overall pattern or syndrome of a strong decline of Pannonian or steppic species in Central Europe following agricultural intensification after the Second World War (Hejda et al. 2017; Kratschmer et al. 2021; Wood and Patiny 2025). Whilst this trend is well-established for bumble bees (*Bombus*), taxonomic impediment has limited our understanding of decline in



**Figure 5.** *Nomiapis valga* (Gerstäcker, 1872) male specimen from Zaragoza, Spain (BBSL740380; BBSL-USDA-ARS); hind leg and metasoma, profile view. Photograph by Skyler Burrows.

European *Nomiapis*; the clarity provided here will hopefully allow a concrete classification of conservation concern to be made.

Finally, Warncke (1976: 111) reports *N. femoralis* ssp. *femoralis* from Croatia (Istria, northern Dalmatia). The location of supporting specimens is unclear. It is not at all clear what these may be, as Istria and Dalmatia have a Mediterranean climate, which does not ecologically match the distributional pattern of *N. femoralis* s. str. Croatia must remain an unconfirmed part of the distribution of *N. femoralis*.

**Distribution.** Germany, Czechia, Slovakia, Hungary, ?Croatia; Serbia, Romania, Ukraine, Russia (European part, Urals, Western Siberia, Eastern Siberia), Kazakhstan, Turkmenistan, Uzbekistan, Tajikistan, Kyrgyzstan, Mongolia, China (Warncke 1976 as *Nomia femoralis* ssp.

femoralis; Ebmer 1988; Astafurova and Pesenko 2006; Pauly 2015) (Fig. 4). The specimen reported from Greece (Rhodes) by Baker (2002: 66) is a female and hence is considered to be a misidentification of *N. caucasica*.

#### Nomiapis valga (Gerstäcker, 1872)

Nomia valga Gerstäcker, 1872: 302, ♂ [Spain: Andalusia, RMNH, neotype by present designation] (Fig. 3)

Material examined. Spain • 1 ♂; Granada, Pantano de Cubillas; 27 May—9 Jun. 1982; R. Leys leg.; RMNH; RMNH. INS.1662584 (neotype by present designation) • 2 ♂; Granada, Pantano de Cubillas; 27 May—9 Jun. 1982; R. Leys leg.; RMNH • 1 ♂, 3 ♀; Granada, Canales; 22 Jul. 1978; R. Leys leg.; RMNH; RMNH.INS.1662575 • 1 ♂; Granada, Cenes de la Vega, Sendero del Río Genil; 750 m a.s.l.; 7 Jun. 2021; T.J. Wood leg.; TJWC • 1 ♂, 2 ♀; Granada, Puente de Ojen; 9 Jul. 1985; R. Leys & P. v. d. Hurk leg.; RMNH; RMNH. INS.1662578 • 4 ♂; Granada, Salobreña; 8 May 1983; W. Perraudin leg.; OÖLM • 1 ♀; Granada, Salobreña; 21 May 1983; W. Perraudin leg.; OÖLM • 1 ♂; Zaragoza, 30 km E; 16 Jun. 1974; P.F. Torchio & E. Ase leg.; BBSL740380; BBSL-USDA-ARS (examined by photograph).

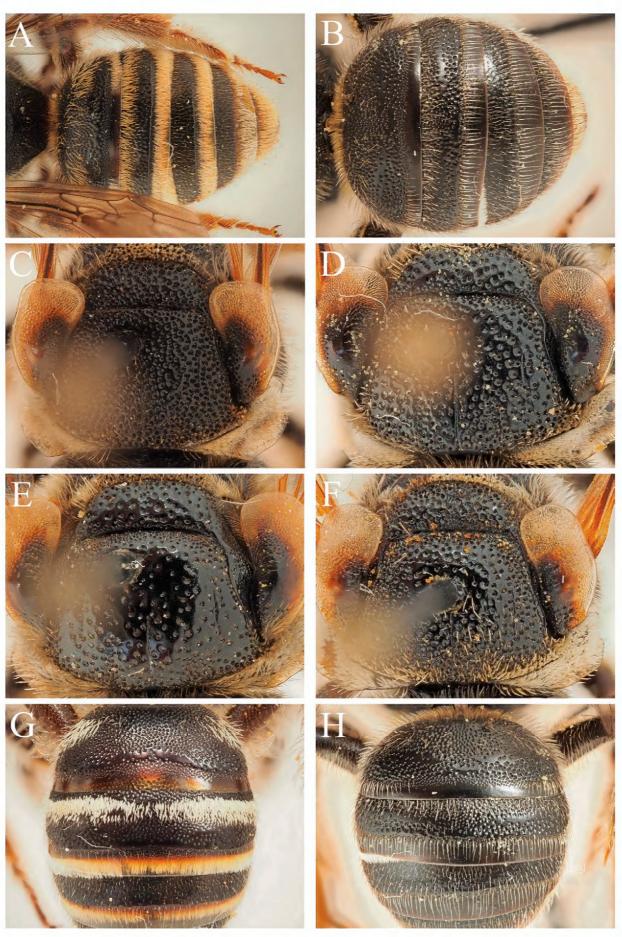
**Notes.** Revision of Spanish material shows that there is a consistent (if infrequently encountered) population in southern Spain, predominantly in the province of Granada but also in Málaga (Pauly 2015); populations in northeastern Spain require investigation, as no records post-1974 are currently available.

**Distribution.** Spain (Málaga, Granada, Zaragoza, ?Barcelona) (Warncke 1976, as *Nomia femoralis* ssp. *valga*; Pauly 2015 *partim*; *present study*) (Fig. 4).

#### Identification key for European *Nomiapis*

Wood and Le Divelec (2022) gave characters to allow separation of members of the *bispinosa* group but not a full key, and currently workers are forced to use either the key of Warncke (1976), which has some problems with precision and employs out-of-date species concepts and nomenclature, or the key of Baker (2002), which deals only with males and also uses some out-of-date species concepts. Both of these keys provide valuable information, but their flaws impede access to the genus for inexperienced workers. Therefore, a new key to the 11 species of *Nomiapis* found in Europe is presented. Only European distributions are given within the identification key, not global distributions, for which broader works should be consulted.

The female of *Nomiapis susannae* Arens, 2018, is currently unknown; it is expected to be very similar to the female of *Nomiapis diversipes*, but sex association must be confirmed using genetic techniques. Greater care should be taken with female specimens due to their inherently more difficult identification; association should be made with concurrently active males, when possible. This is a group that benefits greatly from the consultation of confidently determined or barcoded specimens. Supplementary photographs can be seen in Arens (2018) and Wood and Le Divelec (2022); the figures of Warncke (1976) can be consulted with care and an appreciation of their limitations.



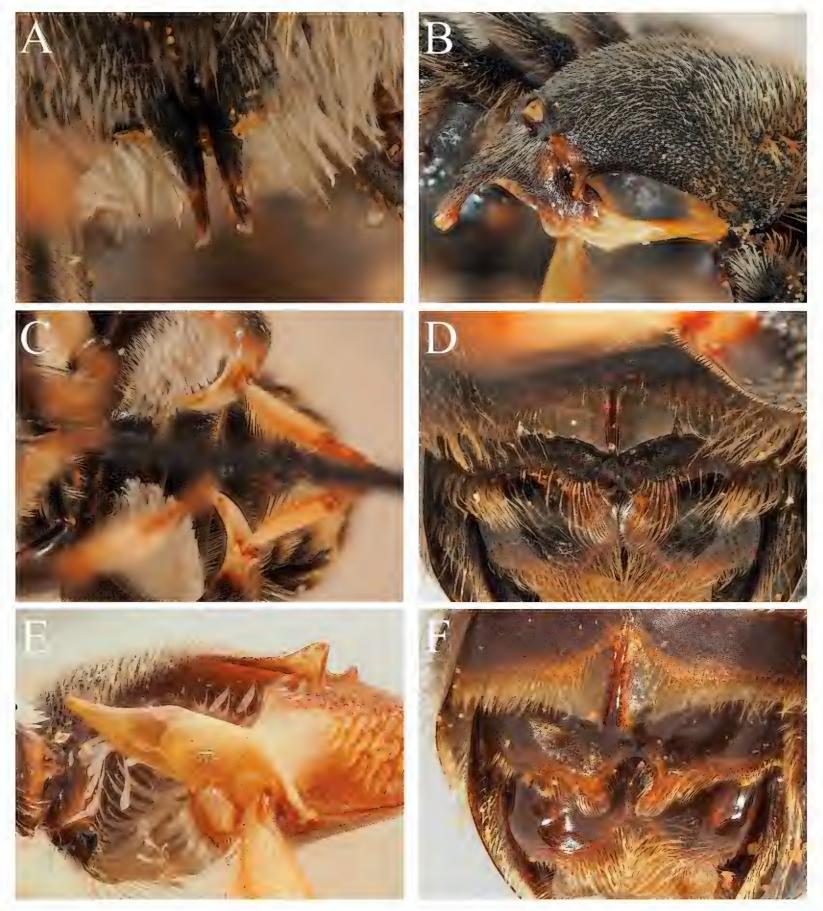
**Figure 6.** *Nomiapis equestris* (Gerstäcker, 1872) female. **A.** Terga, dorsal view. *Nomiapis monstrosa* (Costa, 1861) female; **B.** Terga, dorsal view; **D.** Scutum and scutellum, dorsal view; **H.** T1–2, dorsal view. *Nomiapis diversipes* (Latreille 1806) female; **C.** Scutum and scutellum, dorsal view; **G.** T1–2, dorsal view; *Nomiapis paulyi* Wood & Le Divelec, 2022 female; **F.** Scutum and scutellum, dorsal view.



**Figure 7.** *Nomiapis femoralis* (Pallas, 1773) female. **A.** Vertex, dorsolateral view; **B.** Terga, dorsal view. *Nomiapis valga* (Gerstäcker, 1872) female; **C.** Terga, dorsal view. *Nomiapis caucasica* (Friese, 1897) female; **D.** Terga, dorsal view.

- 8 Head with the posterior margin of the vertex rounded. Found in southern Italy and the Balkan Peninsula.....
- Head with the posterior margin of the vertex displaying a sharp carina (Fig. 7A). Widespread (femoralis-group)....... 9

10 Disc of T1 in basal ½ with punctures extremely dense, separated by < 0.5 puncture diameters, essentially confluent (Fig. 7C). Marginal area of T1 with 5–6 rows of very dense punctures, punctures separated by ≤ 0.5 puncture diameters. Marginal areas of T2-3 with many rows of dense punctures covering their basal ½. Found in Spain (Fig. 4) ...... Disc of T1 in basal  $\frac{1}{2}$  with punctures separated by 0.5–1 puncture diameter (Fig. 7D). Marginal area of T1 with around 4 poorly defined rows of punctures, punctures separated by 0.5–1 puncture diameter. Marginal areas of T2–3 with 1–3 12 Marginal areas of T2–3 with the transition between the disc and the marginal area sharp, the edge almost carina-like. Punctation of marginal areas indistinct, restricted to 2 rows of punctures in basal part. S6 with a pair of strongly produced tubercles medially, clearly forming a strong triangular shape. Slightly larger, 10–12 mm. Distributed across southern Europe in areas with salty soils (coastal habitats, inland saline lagoons, salt steppe) ..... N. bispinosa (Brullé) Marginal areas of T2-3 with the transition between the disc and the marginal area relatively smooth. Punctation of marginal areas distinct, comprising around 4 rows of punctures covering at least the basal 1/5 of marginal area. S6 with a pair of weakly produced tubercles medially, forming a weak triangle. Slightly smaller, 7–9 mm. Distributed in



**Figure 8.** *Nomiapis caucasica* (Friese, 1897) male. **A.** Projecting paired spines on the ventral face of the mesosoma; *Nomiapis monstrosa* (Costa, 1861) male; **B.** Hind leg, femur, and base of tibia, profile view. *Nomiapis diversipes* (Latreille 1806) male; **C.** Hind tibial processes, ventral view; **D.** S4–6, ventral view; *Nomiapis equestris* (Gerstäcker, 1872) male; **E.** Hind tibial process, ventral view; **F.** S4–6, ventral view.

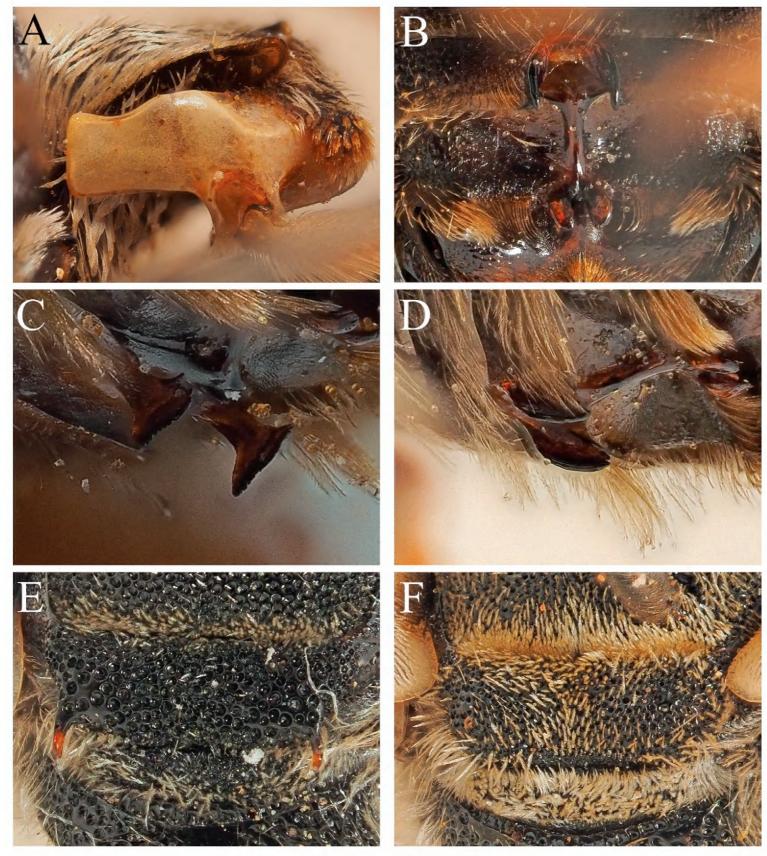
13 Punctures of margins of T2-3 covering basal 1/5 of marginal area. Hind tibia with inner margin comparatively more depressed basally with a subsequent more pronounced angle; inner margin usually crossed over by a sharp carina that Punctures of margins of T2–3 covering slightly more than basal ⅓ of marginal area. Hind tibia with inner margin usually slightly bisinuate, without notable angle following the basal depression; inner margin apically carinate, but the carina usually does not reach the basal depression. Found in Iberia, Corsica, and Sardinia paulyi Wood and Le Divelec 14 Base of the hind tibia with a long digitiform projection (Fig. 8B). Found in southern Italy and the Balkan Peninsula ... N. monstrosa (Costa) 15 Hind tibia produced into a relatively short and apically pointed process (Fig. 8C, E). S4, with its apical margin either straight Hind tibia produced into a relatively long, flattened, and more-or-less rectangular shape with a truncate apex (Fig. 9A). S4, with its apical margin either deeply and broadly excavated medially (Figs 2D, 9B) or, if only narrowly incised, with-Margin of S5 with deep circular emargination, lateral sides produced into curving digiform projections, almost hyaline, 16 S4 medially with raised longitudinal ridges placed very close together, inner margins parallel (Fig. 8D). S5 with 17 median lobes pronounced, comparatively more strongly projecting posteriorly, outer margin therefore somewhat concave. S6 with lateral ridges more strongly developed, standing above the rest of the disc, apically produced into a rounded tooth-like bump. Widespread across Southern and Central Europe, the most commonly encountered Eu-S4 medially with raised longitudinal ridges with their inner margins diverging apically. S5 with medial lobes weakly produced, only slightly projecting posteriorly, outer margin therefore only weakly concave. S6 with lateral ridges only weakly differentiated from surrounding disc, apically disappearing into underlying sculpture. Currently known only 18 Margin of S4 deeply but narrowly incised, lateral margins only slightly thickened. Margin of S5 with a triangular emargination, without projections internally within this emargination. Found only in southern European Russia ..... Margin of S4 with a deep semi-circular emargination (Figs 2D, 9B), lateral margins with elevated ridges, their surface 19 In profile view, lateral margins flanking the emargination of S4 produced elevated projections with triangular to trape-In profile view, lateral margins flanking the emargination of S4 produced into low-lying posteriorly-projecting ridges 20 Disc of T2 uniformly punctate, punctures extending over the entire breadth of the disc, consistently separated by only narrow interspaces with exception of very slightly shining areas mediolaterally, these not exceeding the diameter of a puncture (Fig. 3C). Marginal areas of T2–3 with 3 rows of punctures, these occupying the basal ⅓–⅓ of the marginal areas. In direct comparison, tergal punctation slightly denser and microsculpture slightly stronger, therefore slightly more dull, particularly visible on tergal margins. Scutellum laterally mucronate (Fig. 9E; with two small posteriorly Disc of T2 irregularly punctate, mediolaterally with two shining areas, in size exceeding the diameter of a puncture (Fig. 2C). Marginal areas of T2–3 with 1 row of puncture, sometimes with a very narrow irregular 2<sup>nd</sup> row composed of very small punctures; punctures occupying <1/s of the basal part of the marginal areas. In direct comparison, tergal punctation slightly sparser and microsculpture slightly weaker; therefore, slightly more brightly shining, particularly visible on 

## Discussion

The fortunate retrieval of the remaining type material of *N. caucasica* hidden in the Warncke collection has allowed its identity to be decisively settled in opposition to *N. valga*, which is now considered to be endemic to Spain. That the identity and distribution of what

is ultimately a small number of species (11 in Europe) could be so confused over time (Friese 1897; Warncke 1976; Van der Zanden 1997; Baker 2002; Wood et al. 2020; Wood and Le Divelec 2022) is testament to the high degree of identification difficulty posed by nomine bees, particularly in the female sex. The result also continues to reinforce the importance of the Iberian

<sup>\*</sup> The following two species are more or less impossible to separate in the male sex but can be identified based on the geographic origin and association with females.



**Figure 9.** *Nomiapis caucasica* (Friese, 1897) male. **A.** Hind tibial process, ventral view; **F.** Scutellum, dorsal view. *Nomiapis valga* (Gerstäcker, 1872) male; **B.** S4–5, ventral view; **D.** Medial projections of S4, profile view; **E.** Scutellum, dorsal view. *Nomiapis femoralis* (Pallas, 1773) male; **C.** Medial projections of S4, profile view.

Peninsula as a center for endemic bee diversity both globally and within Europe, a pattern repeatedly borne out during the past decade of taxonomic research on bees (Ghisbain et al. 2023).

Concerning the revisionary work presented here, it is fortunate that the type series of *N. caucasica* was male and that the original description of *N. valga* was also made from the male sex. This is not the case in many other groups that pose identification challenges, particularly in the Indo-Malayan region with genera such as *Maculonomia* Wu, 1982, for which some of the oldest names were described from females such as *Maculonomia elegans* (Smith, 1857), *Maculonomia interrupta* (Cameron, 1904), *Maculonomia leucozonata* (Cameron, 1902), and *Maculonomia terminata* (Smith, 1875) (Pauly 2009). Some genera, such as *Reepenia* Friese, 1909, or *Mellitidia* Guérin-Méneville, 1831,

have never been properly revised, at least in the modern period (Pauly 2009).

Given the fundamental problems of nomine nomenclature and species concepts in the European fauna, it is likely that many changes will be needed in the Indo-Malayan region when revisionary opportunities become available, quite apart from the known problems posed by generic delineation and classification (Bossert et al. 2021; 2024; 2025). However, the results presented here also show that *Nomiapis* females can be consistently identified with care, association with males, integration of genetic barcode data, consideration of biogeography, and careful assessment of puncture density and integumental sculpture. This may offer a path towards revision of groups with many taxa described from females, should these groups display similar species-specific variation as that presented here.

# Acknowledgments

I would like to thank Hugo Gaspar (University of Coimbra, Portugal), Dominique Zimmermann (NHMW), and Romain Le Divelec (University of Mons, Belgium) for helping me trace type material in the ZMHB, Coimbra, and NHMW museums, including the discovery of the loan of material to Diniz in 1965. I thank Romain Le Divelec, Simone Flaminio (UMONS), and Alain Pauly (Tanzania Wildlife Research Institute, Tanzania) for reviewer comments. I also thank Esther Ockermüller and Martin Schwarz (OOLM) and James Hogan (OUMNH) for access to their collections. I thank Skyler Burrows (Utah State University) for sending pictures of N. valga from Spain. Finally, I thank Christophe Praz (University of Neuchâtel, Switzerland) for his company on a collecting trip to northern Greece in June 2024, where *Nomiapis* caucasica was captured; specimens from Greece were collected under permit A.Π. ΥΠΕΝ/ΔΔΔ/49007/1635 from the Ministry of Environment and Energy. This work was supported by the project EPIC—Developing courses for European wild bee identification, EC DG ENV project Contract No.: 090201/2024/927644/SER/ENV.D2.

## References

- Arens W (2018) *Nomiapis susannae* sp. n., eine neue Bienen-Art aus Griechenland (Hymenoptera, Apoidea, Halictidae). Mitteilungen der Münchner Entomologischen Gesellschaft 108: 103–107.
- Ascher JS, Pickering J (2024) Discover Life bee species guide and world checklist (Hymenoptera, Apidae, Anthophila). https://www.discoverlife.org/mp/20q?guide=Apoidea\_species&flags=HAS [accessed 12 October 2024]
- Astafurova YA, Pesenko YA (2006) Bees of the Subfamily Nomiinae (Hymenoptera, Halictidae) in Russia and Adjacent Countries: An Annotated List. Entomological Review 86(1): 74–84. https://doi.org/10.1134/S0013873806010040
- Baker D (2002) On Palaearctic and Oriental species of the genera *Pseudapis* W.F. Kirby, 1900, and *Nomiapis* Cockerell, 1919. Beiträge zur Entomologie 52(1): 1–83. https://doi.org/10.21248/contrib.entomol.52.1.1-83
- Bleidorn C, Henze K (2021) A new primer pair for barcoding of bees (Hymenoptera, Anthophila) without amplifying the orthologous coxA gene of *Wolbachia* bacteria. BMC Research Notes 14(1): 427. https://doi.org/10.1186/s13104-021-05845-9
- Bossert S, Murray EA, Pauly A, Chernyshov K, Brady SG, Danforth BN (2021) Gene Tree Estimation Error with Ultraconserved Elements: An Empirical Study on *Pseudapis* Bees. Systematic Biology 70(4): 803–821. https://doi.org/10.1093/sysbio/syaa097
- Bossert S, Pauly A, Danforth BN, Orr MC, Murray EA (2024) Lessons from assembling UCEs: A comparison of common methods and the case of *Clavinomia* (Halictidae). Molecular Ecology Resources 24(3): e13925. https://doi.org/10.1111/1755-0998.13925
- Bossert S, Freitas FV, Pauly A, Zhu G, Crowder DW, Orr MC, Dorey JB, Murray EA (2025) Phylogeny, antiquity, and niche occupancy of *Trinomia* (Hymenoptera, Halictidae), an Afrotropical endemic

- genus of Nomiinae. Molecular Phylogenetics and Evolution 204: 108273. https://doi.org/10.1016/j.ympev.2024.108273
- De Coster W, D'Hert S, Schultz DT, Cruts M, Van Broeckhoven C (2018) NanoPack: Visualizing and processing long-read sequencing data. Bioinformatics (Oxford, England) 34(15): 2666–2669. https://doi.org/10.1093/bioinformatics/bty149
- Ebmer AW (1988) Kritische Liste der nicht-parasitischen Halictidae Österreichs mit Berücksichtigung aller mitteleuropäischen Arten (Insecta, Hymenoptera, Apoidea: Halictidae). Linzer Biologische Beitrage 20(2): 527–711.
- Ebmer AW (2014) Die nicht-parasitischen Halictidae der Insel Zypern in Vergleich zu Kreta mit einer Monographie der *Lasioglossum bimaculatum*-Artengruppe und einer Ubersicht der *Halictus nic-osiae*-Untergruppe (Insecta, Hymenoptera, Apoidea, Halictidae). Linzer Biologische Beitrage 46(1): 291–413.
- Ewels P, Magnusson M, Lundin S, Käller M (2016) MultiQC:Summarize analysis results for multiple tools and samples in a single report. Bioinformatics (Oxford, England) 32(19): 3047–3048. https://doi.org/10.1093/bioinformatics/btw354
- Friese H (1897) Monographie der Bienengattung *Nomia* (Latr.) (Palaearctische Formen). In: Fest-Schrift zur Feier des fünfzigjahrigen Bestehens des Vereins für schlesische Insektenkunde in Breslau 1897, 45–84.
- Gerstäcker A (1872) Hymenopterische beiträge. Stettiner Entomologische Zeitung 33: 250–308.
- Ghisbain G, Rosa P, Bogusch P, Flaminio S, Le Divelec R, Dorchin A, Kasparek M, Kuhlmann M, Litman J, Mignot M, Müller A, Praz C, Radchenko VG, Rasmont P, Risch S, Roberts SPM, Smit J, Wood TJ, Michez D, Reverté S (2023) The new annotated checklist of the wild bees of Europe (Hymenoptera, Anthophila). Zootaxa 5327(1): 1–147. https://doi.org/10.11646/zootaxa.5327.1.1
- Gouy M, Guindon S, Gascuel O (2010) SeaView version 4: A multiplatform graphical user interface for sequence alignment and phylogenetic tree building. Molecular Biology and Evolution 27(2): 221–224. https://doi.org/10.1093/molbev/msp259
- Hejda R, Farkač J, Chobot K (2017) Červený seznam ohrožených druhů České republiky. Bezobratlí. List of threatened species in the Czech Republic. Invertebrates. Agentura ochrany přírody a krajiny ČR, Praha, 611 pp.
- ICZN [International Commission on Zoological Nomenclature] (1999) International Code of Zoological Nomenclature. 4<sup>th</sup> edn. The International Trust for Zoological Nomenclature, London, xxix + 306 pp.
- Ivanova NV, Dewaard JR, Hebert PDN (2006) An inexpensive, automation-friendly protocol for recovering high-quality DNA. Molecular Ecology Notes 6(4): 998–1002. https://doi.org/10.1111/j.1471-8286.2006.01428.x
- Katoh K, Standley DM (2013) MAFFT multiple sequence alignment software version 7: Improvements in performance and usability. Molecular Biology and Evolution 30(4): 772–780. https://doi.org/10.1093/molbev/mst010
- Kratschmer S, Zettel H, Ockermüller E, Zimmermann D, Schoder S, Neumayer J, Gusenleitner F, Zenz K, Mazzucco K, Ebmer AW, Kuhlmann M (2021) Threat Ahead? An Experts' Opinion on the Need for Red Lists of Bees to Mitigate Accelerating Extinction Risks – The Case of Austria. Bee World 98(3): 74–77. https://doi.or g/10.1080/0005772X.2021.1940734
- Kumar S, Stecher G, Li M, Knyaz C, Tamura K (2018) MEGA X: Molecular Evolutionary Genetics Analysis across computing platforms.

- Molecular Biology and Evolution 35(6): 1547–1549. https://doi.org/10.1093/molbev/msy096
- Michener CD (2007) The Bees of the World. 2<sup>nd</sup> edn. The Johns Hopkins University Press, Baltimore, USA, 953 pp.
- Mölder F, Jablonski KP, Letcher B, Hall MB, Tomkins-Tinch CH, Sochat V, Forster J, Lee S, Twardziok SO, Kanitz A, Wilm A, Holtgrewe M, Rahmann S, Nahnsen S, Köster J (2021) Sustainable data analysis with Snakemake. F1000 Research 10: 33. https://doi.org/10.12688/f1000research.29032.2
- Pallas PS (1773) Reise durch verschiedene Provinzen des Russischen Reichs, zweiter Theil, Erstes Buch vom Jahr 1770. Imperial Academy of Sciences, St. Petersburg, 368 pp.
- Pauly A (1990) Classification des Nomiinae africains (Hymenoptera, Apoidea, Halictidae). Annales Sciences zoologiques, Musée royal de l'Afrique centrale 261: 1–206.
- Pauly A (2009) Classification des Nomiinae de la Région Orientale, de Nouvelle-Guinée et des îles de l'Océan Pacifique (Hymenoptera, Apoidea, Halictidae). Bulletin de l'Institut royal des sciences naturelles de Belgique 79: 151–229.
- Pauly A (2014) Les Abeilles des Graminées ou *Lipotriches* Gerstaecker, 1858, sensu stricto (Hymenoptera, Apoidea, Halictidae, Nomiinae) de l'Afrique subsaharienne. Belgian Journal of Entomology 20: 1–393.
- Pauly A (2015) Le genre *Nomiapis* Cockerell 1919. http://www.atlashymenoptera.net/page.aspx??ID=72 [accessed 30 December 2024]
- Pauly A, Brooks RW, Nilsson LA, Pesenko YA, Eardley CD, Terzo M, Griswold T, Schwarz M, Patiny S, Munzinger J, Barbier Y (2001)

- Hymenoptera Apoidea de Madagascar et des îles voisines. Annales Sciences zoologiques, Musée royal de l'Afrique centrale 286: 1–390.
- Sahlin K, Lim MCW, Prost S (2021) NGSpeciesID: DNA barcode and amplicon consensus generation from long-read sequencing data. Ecology and Evolution 11(3): 1392–1398. https://doi.org/10.1002/ece3.7146
- Van der Zanden G (1997) Einige Notizen über die Gattungen *Pseudapis* Kirby 1900 und *Nomia* Latreille 1804 in der Paläarktis. Linzer Biologische Beitrage 29(2): 753–757.
- Warncke K (1976) Zur Systematik und Verbreitung der Bienengattung *Nomia* Latr. in der Westpalärtis und dem turkestanischen Becken (Hymenoptera, Apoidea). Reichenbachia 16: 93–120.
- Warncke K (1980) Die Bienengattung *Nomia* und *Systropha* im Iran mit Ergänzungen zu den *Nomia*-Arten der Westpaläarktis. Linzer Biologische Beitrage 12(2): 363–384.
- Wood TJ, Le Divelec R (2022) Cryptic Diversity Revealed in A Revision of West Palaearctic *Nomiapis* and *Systropha* (Hymenoptera, Halictidae). Diversity 14(11): 920. https://doi.org/10.3390/d14110920
- Wood TJ, Patiny S (2025) Notes on the genera *Ammobatoides* and *Melitturga* in Europe (Hymenoptera, Apoidea). Journal of Melittology 128(128): 1–21. https://doi.org/10.17161/jom.vi128.23023
- Wood TJ, Cross I, Baldock DW (2020) Updates to the bee fauna of Portugal with the description of three new Iberian *Andrena* species (Hymenoptera, Apoidea, Anthophila). Zootaxa 4790(2): 201–228. https://doi.org/10.11646/zootaxa.4790.2.1